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Leu Pro Pro 35	Gly Pro Let	ı Gly Trp I 40	Pro Leu Val	Gly Ala Leu 45	Pro Leu
Leu Gly Ala 50	Met Pro His	Val Ala I 55	Leu Ala Lys	Leu Ala Lys 60	Lys Tyr
Gly Pro Ile 65	Met His Leu 70	ı Lys Met (Gly Thr Cys 75	Asp Met Val	Val Ala 80
Ser Thr Pro	Glu Ser Ala 85	a Arg Ala I	Phe Leu Lys 90	Thr Leu Asp	Leu Asn 95
Phe Ser Asn	Arg Pro Pro		Gly Ala Ser 105	His Leu Ala 110	Tyr Gly
Ala Gln Asp 115	Leu Val Phe	e Ala Lys 1 120	Tyr Gly Pro	Arg Trp Lys 125	Thr Leu
Arg Lys Leu 130	Ser Asn Leu	ı His Met I 135	Leu Gly Gly	Lys Ala Leu 140	Asp Asp
Trp Ala Asn 145	Val Arg Val		Leu Gly His 155	Met Leu Lys	Ala Met 160
Cys Glu Ala	Ser Arg Cys	s Gly Glu I	Pro Val Val 170	Leu Ala Glu	Met Leu 175

Thr Tyr Ala Met Ala Asn Met Ile Gly Gln Val Ile Leu Ser Arg Arg 180 185 190

tgggtatttt atccatttaa tttttcttca gaattatgat catagttatc agaataaaat

1740

Val Phe Val Thr Lys Gly Thr Glu Ser Asn Glu Phe Lys Asp Met Val 195 200 205

Val Glu Leu Met Thr Ser Ala Gly Tyr Phe Asn Ile Gly Asp Phe Ile 210 215 220

Pro Ser Ile Ala Trp Met Asp Leu Gln Gly Ile Glu Arg Gly Met Lys 225 230 235 240

Lys Leu His Thr Lys Phe Asp Val Leu Leu Thr Lys Met Val Lys Glu 245 250 255

His Arg Ala Thr Ser His Glu Arg Lys Gly Lys Ala Asp Phe Leu Asp 260 265 270

Val Leu Leu Glu Glu Cys Asp Asn Thr Asn Gly Glu Lys Leu Ser Ile 275 280 285

Thr Asn Ile Lys Ala Val Leu Leu Asn Leu Phe Thr Ala Gly Thr Asp 290 295 300

Thr Ser Ser Ser Ile Ile Glu Trp Ala Leu Thr Glu Met Ile Lys Asn 305 310 315 320

Pro Thr Ile Leu Lys Lys Ala Gln Glu Glu Met Asp Arg Val Ile Gly 325 330 335

Arg Asp Arg Leu Leu Glu Ser Asp Ile Ser Ser Leu Pro Tyr Leu 340 345 350

Gln Ala Ile Ala Lys Glu Thr Tyr Arg Lys His Pro Ser Thr Pro Leu 355 360 365

Asn Leu Pro Arg Ile Ala Ile Gln Ala Cys Glu Val Asp Gly Tyr Tyr 370 375 380

Ile Pro Lys Asp Ala Arg Leu Ser Val Asn Ile Trp Ala Ile Gly Arg 385 390 395 400

Asp Pro Asn Val Trp Glu Asn Pro Leu Glu Phe Leu Pro Glu Arg Phe
405 410 415

Leu Ser Glu Glu Asn Gly Lys Ile Asn Pro Gly Gly Asn Asp Phe Lys

420 425 430

Leu Ile Pro Phe Gly Ala Gly Arg Arg Ile Cys Ala Gly Thr Arg Met 435 440 445

Gly Met Val Leu Val Ser Tyr Ile Leu Gly Thr Leu Val His Ser Phe 450 455 460

Asp Trp Lys Leu Pro Asn Gly Val Ala Glu Leu Asn Met Asp Glu Ser 465 470 475 480

Phe Gly Leu Ala Leu Gln Lys Ala Val Pro Leu Ser Ala Leu Val Ser 485 490 495

Pro Arg Leu Ala Ser Asn Pro Tyr Ala Thr

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<213> salvia

<400> 13

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tattaaagcc cttttactga atttatttac tqcaqqqaca qacacatcqt cqaqcataat 960 agaatgggca ttagcggaga tgatcaagaa tccaagcatc caaaaaaggg ctcaccaaga 1020 gatggacaga gtcatcggga gagagcggcg tttgctcgaa tccgacatcc caaatctqcc 1080 atacctcaaa gccatatgca aagaggcata ccgaaaacac ccttccacgc cactaaacct 1140 gcctcggatc tccacggatg catgcgtcgt cgatggctac cacatcccca agaacacgag 1200 gttgagcgtc aacatctggg ccataggccg agatcccgac gtttgggaga atccccttga 1260 cttcaaccct gacaggttta tgtcagggtt gcaggggatt gagcccggag ggaatcactt 1320 cgagctcatt ccctttgggg cggggcgcag gatctgcgcc ggcagcagaa tggggattgt 1380 aatagtggag tatttgctgg cgacactcgt gcactctttc gaatgggatt tgccggccgg 1440 ctcagcggag atggacatgg aggaggtgtt cggqctqqcc ttqcaqaaaq ctqtaccact 1500 tgctgctagg ctcactccta ggttgccttc acattgctat gcacctcctt ctatttaatt 1560 tgcatattta catatgttgt gttacattga gcctttqcat atqttqtatc caacctatct 1620 tataacttgt gcatgaaatt gaaaaaaaaa aaaaaaaaa 1659

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<211> 520

<212> PRT

<213> salvia

<400> 14

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Lys Leu Ser Thr Thr Gly His Pro Leu Pro Pro Gly Pro Arg Gly Phe 35 40 45

Leu Val Val Gly Ser Leu Pro Leu Leu Gly Asp Met Pro His Val Ala 50 55 60

Leu Ala Lys Met Ala Lys Thr Tyr Gly Pro Ile Met Tyr Leu Lys Met 65 70 75 80

Gly Thr Val Gly Met Val Val Ala Ser Thr Pro Asp Ala Ala Arg Ala 85 90 95

Phe Leu Lys Thr His Asp Ala Asn Phe Ser Asn Arg Pro Val Asn Ala

100 105 110

Gly Ala Thr Ile Leu Ala Tyr Asn Ala Gln Asp Met Val Phe Ala Pro 115 120 125

Tyr Gly Pro Lys Trp Arg Leu Leu Arg Lys Leu Ser Ser Leu His Met 130 135 140

Leu Gly Ser Lys Ala Leu Glu Glu Trp Ala Asp Val Arg Thr Ser Glu
145 150 155 160

Val Gly His Met Leu Ala Ala Met His Glu Ala Ser Arg Leu Gly Glu 165 170 175

Ala Val Gly Leu Pro Glu Met Leu Val Tyr Ala Thr Ala Asn Met Ile 180 185 190

Gly Gln Val Ile Leu Ser Arg Arg Val Phe Val Thr Lys Gly Lys Glu 195 200 205

Met Asn Glu Phe Lys Glu Met Val Val Glu Leu Met Thr Thr Ala Gly 210 215 220

Tyr Phe Asn Ile Gly Asp Phe Ile Pro Trp Leu Ala Trp Met Asp Leu 225 230 235 240

Gln Gly Ile Glu Arg Gly Met Lys Lys Leu His Lys Lys Trp Asp Arg 245 250 255

Leu Ile Gly Lys Met Leu Asp Asp Arg Leu Lys Ser Thr Tyr Lys Arg 260 265 270

Asn Asp Lys Pro Asp Leu Leu Asp Ser Leu Leu Ala Asn His Asp Asp 275 280 285

Glu Ser Lys Asp Asp Asp Glu Asp Cys Lys Leu Thr Thr Thr Asn Ile 290 295 300

Lys Ala Leu Leu Leu Asn Leu Phe Thr Ala Gly Thr Asp Thr Ser Ser 305 310 315 320

Ser Ile Ile Glu Trp Ala Leu Ala Glu Met Ile Lys Asn Pro Ser Ile 325 330 335

	340		;	345				350		
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Cys Lys Glu 370	Ala Tyr	Arg Lys 375	His 1	Pro Ser	Thr	Pro 380	Leu	Asn	Leu	Pro
Arg Ile Ser 385	Thr Asp	Ala Cys 390	Val V	Val Asp	Gly 395	Tyr	His	Ile	Pro	Lys 400
Asn Thr Arg	Leu Ser 405	Val Asn	Ile :	Trp Ala 410	Ile	Gly	Arg	Asp	Pro 415	Asp
Val Trp Glu	Asn Pro 420	Leu Asp		Asn Pro 425	Asp	Arg	Phe	Met 430	Ser	Gly
Leu Gln Gly 435	Ile Glu	Pro Gly	Gly 1 440	Asn His	Phe	Glu	Leu 445	Ile	Pro	Phe
Gly Ala Gly 450	Arg Arg	Ile Cys 455	Ala (Gly Ser	Arg	Met 460	Gly	Ile	Val	Ile
Val Glu Tyr 465	Leu Leu	Ala Thr 470	Leu V	Val His	Ser 475	Phe	Glu	Trp	Asp	Leu 480
Pro Ala Gly	Ser Ala 485	Glu Met	Asp I	Met Glu 490	Glu	Val	Phe	Gly	Leu 495	Ala
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Gln Lys Arg Ala His Gln Glu Met Asp Arg Val Ile Gly Arg Glu Arg

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<210> 16 <211> 518 <212> PRT <213> salvia

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1

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35 40 45

Gly Ser Leu Pro Leu Leu Gly Asp Met Pro His Val Ala Leu Ala Lys 50 55 60

Met Ala Lys Thr Tyr Gly Pro Ile Met Tyr Leu Lys Met Gly Thr Val 65 70 75 80

Gly Met Val Val Ala Ser Thr Pro Asp Ala Ala Arg Ala Phe Leu Lys 85 90 95

Thr Gln Asp Ala Asn Phe Ser Asn Arg Pro Val Asn Ala Gly Ala Thr 100 105 110

Ile Leu Ala Tyr Asn Ala Gln Asp Met Val Phe Ala Pro Tyr Gly Pro 115 120 125

Lys Trp Arg Leu Leu Arg Lys Leu Ser Ser Leu His Met Leu Gly Ser 130 135 140

Lys Ala Leu Glu Glu Trp Ala Asp Val Arg Thr Ser Glu Val Gly His 145 150 155 160

Met Leu Ala Ala Met His Glu Ala Ser Arg Leu Asp Glu Ala Val Gly
165 170 175

Leu Pro Glu Met Leu Val Tyr Ala Thr Ala Asn Met Ile Gly Lys Val

Ile Leu Ser Arg Arg Val Phe Val Thr Lys Gly Lys Glu Met Asn Glu 195 200 205

Phe Lys Glu Met Val Val Glu Leu Met Thr Thr Ala Gly Tyr Phe Asn 210 220

Ile Gly Asp Phe Ile Pro Trp Leu Ala Trp Met Asp Leu Gln Gly Ile 225 230 235 240

Glu Arg Gly Met Lys Lys Leu His Lys Lys Trp Asp Arg Leu Ile Gly 245 250 Lys Met Leu Asp Asp Arg Leu Lys Ser Thr Tyr Lys Arg Asn Asp Lys 260 265 Pro Asp Leu Leu Asp Ser Leu Leu Ala Asn His Asp Asp Glu Ser Lys Asp Asp Glu Asp Cys Lys Leu Thr Thr Asn Ile Lys Ala Leu 295 Leu Leu Asn Leu Phe Thr Ala Gly Thr Asp Thr Ser Ser Ser Ile Ile Glu Trp Ala Leu Ala Glu Met Ile Lys Asn Pro Ser Ile Gln Lys Arg Ala His Gln Glu Met Asp Arg Val Ile Gly Arg Glu Arg Arg Leu Leu Glu Ser Asp Ile Pro Asn Leu Pro Tyr Leu Lys Ala Ile Cys Lys Glu Ala Tyr Arg Lys His Pro Ser Thr Pro Leu Asn Leu Pro Arg Ile Ser 370 375 Thr Asp Ala Cys Val Val Asp Gly Tyr His Ile Pro Lys Asn Thr Arq 390 Leu Ser Val Asn Ile Trp Ala Ile Gly Arg Asp Pro Asp Val Trp Glu 405 410 Asn Pro Leu Asp Phe Asn Pro Asp Arg Phe Met Ser Gly Leu Gln Gly 420 425 Ile Glu Pro Gly Gly Asn His Phe Glu Leu Ile Pro Phe Gly Ala Gly Arg Arg Ile Cys Ala Gly Ser Arg Met Gly Ile Val Ile Val Glu Tyr 455

Leu Leu Ala Thr Leu Val His Ser Phe Glu Trp Asp Leu Pro Ala Gly

470

Ser Ala Glu Met Asp Met Glu Glu Val Phe Gly Leu Ala Leu Gln Lys 485

Ala Val Pro Leu Ala Ala Arg Leu Thr Pro Arg Leu Pro Ser His Cys 505

Tyr Ala Pro Pro Ser Ile 515

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<212> DNA

<213> sollya

<220>

<221> misc feature

<222> (1372)..(1372)

<223> n = any nucleotide

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<212> PRT

<213> sollya

<400> 18

Met Ala Thr Thr Leu Glu Phe Ile Leu Cys Phe Thr Ile Thr Ala Leu 1 5 10 15

Pro Phe Leu Tyr Cys Ile Leu Asn Met Arg Ile Leu Leu Asn Arg His 20 25 30

Pro Arg Ser Leu Pro Pro Gly Pro Arg Pro Trp Pro Ile Val Gly Asn 35 40 45

Leu Pro His Leu Gly Thr Lys Pro His His Ser Ile Ala Ala Met Ala 50 55 60

Arg Lys Tyr Gly Pro Leu Leu His Leu Arg Met Gly Ile Val His Val 65 70 75 80

Val Val Ala Ala Ser Ala Asp Val Ala Ala Gln Phe Leu Lys Asn Asp 85 90 95

Ala Asn Phe Ser Ser Arg Pro Pro Asn Ser Gly Ala Lys His Met Ala 100 105 110

Tyr Asn Tyr His Asp Met Val Phe Ala Pro Tyr Gly Pro Arg Trp Arg Met Leu Arg Lys Ile Cys Ala Leu His Ile Phe Ser Ala Lys Ala Leu Asp Asp Phe His Arg Val Arg Glu Glu Val Ala Ile Leu Ala Arg Thr Leu Ala His Ala Gly Gln Lys Pro Val Asn Leu Gly Gln Leu Phe Ser Thr Cys Asn Ala Asn Ala Leu Ser Val Leu Met Leu Gly Arg Arg Leu Phe Ser Thr Glu Val Asp Ser Lys Ala Tyr Asp Phe Lys Gln Met Val Val Glu Leu Met Thr Leu Ala Gly Glu Phe Asn Val Ser Asp Phe Ile Pro Pro Leu Glu Trp Leu Asp Leu Gln Gly Val Ala Ala Lys Met Lys Asn Val His Asn Arg Phe Asp Ala Phe Leu Asn Val Ile Leu Glu Glu His Lys Leu Lys Leu Asn Asn Ser Gly His Gly Glu Gln Lys His Met Asp Leu Leu Ser Thr Leu Ile Leu Lys Asp Asp Ala Asp Ser Glu Gly Gly Lys Leu Thr Asp Thr Glu Ile Lys Ala Leu Leu Leu Asn Leu Phe Ser Ala Gly Thr Asp Thr Ser Ser Ser Thr Ile Glu Trp Val Ile Ala Glu Leu Ile Arg Asn Pro Lys Ile Leu Ala Gln Ala Gln Arg

Glu Leu Asp Leu Val Val Gly Pro Asn Arg Leu Val Thr Asp Leu Asp

340 345 350

Leu Lys Gln Leu Thr Tyr Leu Gln Ala Ile Val Lys Glu Thr Phe Arg 355 360 365

Leu His Pro Ala Thr Pro Leu Ser Leu Pro Arg Ile Ala Thr Glu Ser 370 375 380

Cys Glu Ile Asn Gly Phe Tyr Ile Pro Lys Gly Ser Thr Leu Leu Val 385 390 395 400

Asn Ile Trp Ala Ile Gly Arg Asp Pro Asn Thr Trp Ala Glu Pro Leu
405 410 415

Val Phe Arg Pro Glu Arg Phe Leu Ser Asp Gly Glu Ser Pro Asn Val 420 425 430

Asp Val Lys Gly Arg Asn Phe Glu Leu Ile Pro Phe Gly Ala Gly Arg 435 440 445

Arg Ile Cys Ala Gly Met Asn Phe Gly Leu Arg Met Val Gln Leu Val 450 455 460

Thr Ala Thr Leu Ile His Ala Phe Asn Trp Glu Leu Pro Glu Gly Glu 465 470 475 480

Leu Pro Glu Asn Met Asn Met Glu Glu Asp Tyr Gly Ile Ser Leu Gln
485 490 495

Arg Thr Val Pro Leu Val Val His Pro Lys Pro Arg Leu Asp His Glu 500 505 510

Val Tyr Gln Ser His Gly Val Val Asn 515 520

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<211> 37

<212> DNA

<213> petunia

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Arg	Lys 130	Leu	Ser	Asn	Leu	His 135	Met	Leu	Gly	Gly	Lys 140	Ala	Leu	Glu	Glu
Trp 145	Ser	Gln	Val	Arg	Glu 150	Ile	Glu	Met	Gly	His 155	Met	Leu	Arg	Ala	Met 160
Tyr	Asp	Cys	Ser	Gly 165	Gly	Gly	Asp	Gly	Asn 170	Asn	Asp	Asn	Asp	Gly 175	Asn
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Cys Ile Pro Arg Gln Arg Val Arg Asn Arg Thr Ser Leu Arg Thr Trp 210 215 220

Trp Phe Lys Leu Met Thr Val Ala Gly Tyr Phe Asn Ile Gly Asp Phe 225 230 235 240

Phe Pro Phe Leu Ala Arg Arg Arg Gln Gly Ile Glu Arg Gly Met 245 250 255

Lys Thr Leu His Asn Lys Lys Asp Asp Leu Leu Thr Thr Met Ile His 260 265 270

Glu His Val Ala Ser Ala His Lys Arg Lys Gly Lys Pro Pro Phe Leu 275 280 285

Asp Val Leu Met Ala His His Thr Asn Glu Ser His Glu Leu Ser Leu 290 295 300

Thr Asn Ile Lys Ala Leu Leu Leu Asn Leu Phe Thr Ala Gly Thr Asp 305 310 315 320

Thr Ser Ser Ser Ile Ile Glu Trp Ala Leu Ala Glu Met Leu Ile Asn 325 330 335

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Gln Ala Ile Cys Lys Glu Thr Tyr Arg Lys His Pro Ser Thr Pro Leu 370 375 380

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Asp Pro Asn Val Trp Glu Asn Pro Leu Glu Phe Asn Pro Glu Arg Phe

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±20	425	430

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Pro Ile Leu Gly Ala Leu Pro Leu Gly Asn Met Pro His Val Thr 50 55 60

Phe Ala Asn Met Ala Lys Lys Tyr Gly Ser Val Met Tyr Leu Lys Val 65 70 75 80

Gly Ser His Gly Leu Ala Ile Ala Ser Thr Pro Asp Ala Ala Lys Ala Phe Leu Lys Thr Leu Asp Leu Asn Phe Ser Asn Arg Pro Pro Asn Ala 105 Gly Ala Thr His Leu Ala Tyr Asn Ala Gln Asp Met Val Phe Ala His 120 Tyr Gly Pro Lys Trp Lys Leu Leu Arg Lys Leu Ser Asn Leu His Met 135 Leu Gly Gly Lys Ala Leu Glu Asn Trp Ala Asp Val Arg Lys Thr Glu 150 Leu Gly Tyr Met Leu Lys Ala Met Phe Glu Ser Ser Gln Asn Asn Glu 165 170 Pro Val Met Ile Ser Glu Met Leu Thr Tyr Ala Met Ala Asn Met Leu 180 185 Ser Gln Val Ile Leu Ser Arg Arg Val Phe Asn Lys Lys Gly Ala Lys 195 200 Ser Asn Glu Phe Lys Asp Met Val Val Glu Leu Met Thr Ser Ala Gly 210 215 220 Tyr Phe Asn Ile Gly Asp Phe Ile Pro Ser Ile Gly Trp Met Asp Leu 225 230 235 Gln Gly Ile Glu Gly Gly Met Lys Arg Leu His Lys Lys Phe Asp Val 245 Leu Leu Thr Arg Leu Leu Asp Asp His Lys Arg Thr Ser Gln Glu Arg 260 Lys Gln Lys Pro Asp Phe Leu Asp Phe Val Ile Ala Asn Gly Asp Asn 275 Ser Asp Gly Glu Arg Leu Asn Thr Asp Asn Ile Lys Ala Leu Leu Leu 290 Asn Leu Phe Thr Ala Gly Thr Asp Thr Ser Ser Ser Ile Ile Glu Trp 305

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Asp Glu Met Asp Arg Val Ile Gly Arg Asp Arg Arg Leu Leu Glu Ser 340 345 350

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Asn Pro Asn Glu Phe Asp Pro Glu Arg Phe Leu Tyr Gly Arg Asn Ala 420 425 430

Lys Ile Asp Pro Arg Gly Asn His Phe Glu Leu Ile Pro Phe Gly Ala 435 440 445

Gly Arg Arg Ile Cys Ala Gly Thr Arg Met Gly Ile Leu Leu Val Glu 450 460

Tyr Ile Leu Gly Thr Leu Val His Ser Phe Asp Trp Lys Leu Gly Phe 465 470 475 480

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Gly Ala Leu Pro Val Leu Gly Ser Met Pro His Val Thr Leu Ser Arg 50 55 60

Met Ala Lys Lys Tyr Gly Pro Val Met Tyr Leu Lys Met Gly Thr Lys 65 70 75 80

Asn Met Val Val Ala Ser Thr Pro Ala Ala Ala Arg Ala Phe Leu Lys 85 90 95

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<212> PRT

<213> chrysanthemum

<400> 29

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Ile Asp Gly His Leu Arg Glu Val Gly Leu Thr Phe His Leu Leu Lys

265

245

260

Asp	Val	Pro 275	Gly	Leu	Ile	Ser	Lys 280	Asn	Ile	Glu	Lys	Ala 285	Leu	Thr	Gln		
Ala	Phe 290	Ser	Pro	Leu	Gly	Ile 295	Ser	Asp	Trp	Asn	Ser 300	Ile	Phe	Trp	Ile		
Ala 305	His	Pro	Gly	Gly	Pro 310	Ala	Ile	Leu	Asp	Gln 315	Val	Glu	Leu	Lys	Leu 320		
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Arg	Lys	Lys 355	Ser	Ala	Glu	Glu	Gly 360	Ala	Ala	Thr	Thr	Gly 365	Glu	Gly	Leu		
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ggc	catg	att a	actt	agta	cc a	catg	taac	t ga	gact	tgca	atg	gaca	agt .	actt	attatc	18	30
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<212> PRT

<213> lavendula

<400> 32

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Thr Thr Ile Thr His Gly Gly Ser His Arg Leu Pro Pro Gly Pro Arg
35 40 45

Gly Phe Pro Ile Val Gly Ala Leu Pro Leu Leu Gly Asp Met Pro His 50 55 60 Val Ala Leu Ala Lys Met Ala Lys Thr Tyr Gly Pro Ile Ile Tyr Leu 65 70 75 80

Lys Val Gly Ala Trp Gly Met Ala Val Ala Ser Thr Pro Ala Ser Ala 85 90 95

Arg Ala Phe Leu Lys Thr Leu Asp Thr Asn Phe Ser Asp Arg Pro Pro 100 105 110

Asn Ala Gly Ala Thr Ile Leu Ala Tyr Asn Ala Glu Asp Met Val Phe 115 120 125

Ala Arg Tyr Gly Pro Lys Trp Arg Leu Leu Arg Lys Leu Thr Asn Leu 130 135 140

His Met Leu Gly Asn His Ala Leu Asp Gly Trp Ala Ser Val Arg Ser 145 150 155 160

Ser Glu Leu Gly Tyr Met Leu His Ala Arg His Asp Ala Thr Arg His 165 170 175

Gly Glu Pro Val Val Leu Pro Glu Met Leu Met Tyr Ala Val Gly Asn 180 185 190

Met Leu Gly Gln Val Ile Leu Ser Arg Ile Phe Glu Lys Lys Gly 195 200 205

Lys Glu Val Asn Glu Leu Lys Asp Met Val Val Glu Leu Met Thr Ser 210 215 220

Ala Gly Tyr Phe Asn Ile Gly Asp Phe Ile Pro Trp Leu Ala Trp Met 225 230 235 240

Asp Leu Gln Gly Ile Glu Ser Gly Met Lys Lys Leu His Asn Lys Phe 245 250 255

Asp Lys Leu Ile Gly Lys Met Ile Glu Asp His Leu Lys Ser Ala His
260 265 270

Ile Arg Lys Ala Lys Pro Asp Leu Leu Asp Cys Leu Leu Ala Asn Arg 275 280 285

Asp Ser Ser Asp Ala Glu Lys Leu Thr Ser Thr Asn Val Lys Ala Leu 290 295 300

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Leu Leu Asn Leu Phe Thr Ala Gly Thr Asp Thr Ser Ser Ser Ile Ile